

## Adoptive transfer of T cells transduced with a chimeric antigen receptor to treat relapsed or refractory acute leukemia: efficacy and feasibility of immunotherapy approaches

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Treatment outcomes of acute leukemia (AL) have not improved over the past several decades and relapse rates remain high despite the availability of aggressive therapies. Conventional relapsed leukemia treatment includes second allogeneic hematopoietic stem cell transplantation (allo-HSCT) and donor lymphocyte infusion (DLI), which in most cases mediate, at best, a modest graft-versus-leukemia effect, although their clinical efficacy is still limited. Although allo-HSCT following myeloablative conditioning is a curative treatment option for younger patients with acute myeloid leukemia (AML) in a first complete remission (CR), allo-HSCT as a clinical treatment is usually limited because of treatment-related toxicity. The overall DLI remission rate is only 15%–42% and 2-year overall survival (OS) is approximately 15%–20%, with a high (40%–60%) incidence of DLI-related graft-versus-host disease (GVHD). Therefore, development of new, targeted treatment strategies for relapsed and refractory AL patients is ongoing. Adoptive transfer of T cells with genetically engineered chimeric antigen receptors (CARs) is an encouraging approach for treating hematological malignancies. These T cells are capable of selectively recognizing tumor-associated antigens and may overcome many limitations of conventional therapies, inducing remission in patients with chemotherapy-refractory or relapsed AL. In this review, we aimed to highlight the current understanding of this promising treatment modality, discussing its adverse effects and efficacy.

**chimeric antigen receptor, acute leukemia, efficacy, feasibility**

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### INTRODUCTION

Allo-HSCT and chemotherapy have been used to successfully treat patients with acute lymphoblastic leukemia (ALL), but the prognosis of adult patients with relapsed ALL is still very poor. Long-term survival rate of adult patients with relapsed ALL depends on the achievement of CR, induced through chemotherapy, followed by allo-HSCT (Fielding et al., 2007; Gokbuget et al., 2012). Unfortunately, many patients will not receive allo-HSCT because of a fail-

ure in achieving second CR following salvage chemotherapy. For patients with AML, the probability of being cured with chemotherapy alone is significantly lower than combined treatment, as only 50%–65% patients achieve long-term survival (Gibson et al., 2005; Lange et al., 2008; Creutzig et al., 2008). Moreover, the prognosis is even worse in elderly patients (Kaspers et al., 2006; Appelbaum et al., 2006). To address the problem of limited therapeutic success of treating relapsed/refractory ALL or AML, adoptive therapy with chimeric antigen receptor expressing T (CAR-T) cells is a promising approach for hematologic malignancies.

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## CAR: GENETICALLY MODIFIED ADOPTIVE CELL THERAPY

CAR-T cells were first described as an adoptive cell therapy in 1989 by Gross and colleagues (Gross et al., 1989). CARs are composed of an extracellular domain that recognizes tumor surface antigens and is linked to an intracellular signaling domain via trans-membrane domains. The extracellular domain consists of antigen-binding variable regions from monoclonal antibody heavy and light chains that are fused into a single protein known as the single-chain variable fragment (Dotti et al., 2014; Jena et al., 2010). First-generation CAR contained a single-signaling domain derived from either TCR- $\zeta$  or FcR- $\gamma$  chain (Eshhar et al., 1993; Moritz et al., 1994). Second-generation CAR included a single co-stimulatory domain derived from either 4-1BB or CD28. Third-generation CAR had two co-stimulatory domains, namely 4-1BB and CD28, and other co-stimulatory molecules (Goker et al., 2016). T cell genetic modification with CAR has several advantages in leukemia treatment, e.g., CAR use is applicable to a broad range of patients irrespective of the HLA phenotype. CAR-T cell approach also overcomes ability of the tumor to escape immunosurveillance by down-regulating HLA molecules on cell surface. Targeting of tumor antigens with CAR-T cells is applicable to any tumor antigens, including glycolipids, proteins, and carbohydrates. CAR-T cell clinical trials are currently conducted.

## ONGOING TRIALS OF TREATING ALL PATIENTS WITH CAR-T CELLS

Until now, 30 CAR-T AL trials have been published. Some of these focused on ALL, targeting CD19 or CD20, whereas others focused on AML, targeting Lewis-Y antigen or CD33. The results are summarized in Table 1.

## PUBLISHED CLINICAL OUTCOMES OF ADOPTIVE THERAPY WITH CAR-MODIFIED T CELLS FOR ALL

Relapsed ALL is difficult to treat despite the availability of aggressive therapies. Promising results have been obtained in recent studies on CD19/CD20-targeted CAR-T (CD19/CD20-CAR-T) for ALL treatment (Kochenderfer et al., 2012). Maude and colleagues conducted research into infusing autologous T cells transduced with CD19-targeted CAR (CD19-CAR-T) in patients with relapsed or refractory ALL. Patients were monitored for clinical response and toxic effects. The clinical results revealed that CR was achieved in 27 (90%) out of 30 patients (children and adults). Sustained remission was achieved, with 67% 6-month event-free survival rate and 78% OS rate. Cytokine-release syndrome (CRS), which later developed in 27% patients, was associated with a higher disease burden.

Davila and colleagues examined 16 patients with relapsed or refractory B cell acute lymphoblastic leukemia (B-ALL) who were treated with CD19-CAR-T. Clinical results revealed 88% overall complete response rate. This therapy was as effective in treating high-risk patients with Philadelphia chromosome-positive Ph<sup>+</sup> disease, as in treating those with relapsed disease after previous allo-HSCT (Davila et al., 2014). Shortly afterwards, Lee and colleagues published outcomes of a clinical investigation concerning 21 patients with relapsed or refractory ALL (including eight who had previously undergone allo-HSCT) who were infused with CD19-CAR-T. Safety of the treatment was evaluated and any toxic effects were fully reversible, with the highest severity, Grade 4, CRS observed in 3 (14%) out of 21 patients (95% CI 3.0–36.3) (Lee et al., 2015). Thus, CD19-CAR-T therapy is feasible, safe, and mediates a potent anti-leukemic effect in patients with ALL.

## PUBLISHED CLINICAL OUTCOMES OF ADOPTIVE THERAPY WITH CAR-MODIFIED T CELLS FOR AML

Despite the improvement in allo-HSCT and chemotherapy-based treatment of AML, the majority of standard or high-risk AML patients will die or relapse. AML is sensitive to T-cell-mediated control in the allo-HSCT setting. Nevertheless, therapeutic approaches directed at inducing autologous T-cell responses in patients have shown limited efficacy (Wen et al., 2002; Kershaw et al., 2006). Therefore, the development of novel therapeutics for treating AML is crucial. Although CD19-CAR-T approaches are proving highly effective against hematologic malignancies (Brenner et al., 2013), a concern has been raised that extending this success to other tumors may be difficult. CD19 antigen (Jensen et al., 2010) is generally restricted to the B cell lineage. In contrast, LeY antigen is a difucosylated carbohydrate antigen widely expressed by AML cells, but with a limited expression in healthy tissues. However, its role and significance for the survival of leukemia cells remain to be elucidated. In 2013, a phase I clinical trial was conducted at the Ludwig Institute for Cancer Research (Ritchie et al., 2013) examining autologous CAR-anti-LeY-T therapy for AML. Safety and post-infusion persistence of adoptively transferred T cells. Neither Grade 3 nor Grade 4 toxicities were observed. One patient achieved a cytogenetic remission whereas in another, with active leukemia, reduction in peripheral blood (PB) blasts was observed. This clinical outcome verified the safety and feasibility of CAR-T therapy in high-risk AML patients and demonstrated durable in vivo persistence. CD123, a trans-membrane chain of interleukin (IL)-3 receptor, is expressed in the majority of AML cells and in many normal cells. Recent preclinical reports have demonstrated that CAR-T cells have the potential to effectively and durably eradicate primitive myeloid blast cells (Mardiros et al., 2013). Mardiros and colleagues con-

**Table 1** Summary of ongoing clinical trials of autologous/allogeneic CAR-T cell treatments for AL

Disease	Target antigen	Gene transfer	CAR signaling domain	Sponsor	Clinical Trial.gov ID	Status
ALL	CD20	Retrovirus	scFv-(4-1BB-CD3 $\zeta$ )	Chinese PLA General Hospital	NCT01735604	Recruiting
ALL	CD19	Lentivirus	scFv-(4-1BB-CD3 $\zeta$ )	Penn Medicine's Abramson Cancer Center	NCT00891215	Completed
ALL	CD19	Lentivirus	scFv-(4-1BB-CD3 $\zeta$ )	University of Pennsylvania	NCT01551043	Completed
ALL	CD19	Retrovirus	scFv-(CD28-CD3 $\zeta$ )	Baylor College of Medicine	NCT00608270	Recruiting
ALL	CD19	Retrovirus	scFv-(CD28-CD3 $\zeta$ )	Memorial Sloan Kettering Cancer Center	NCT01416974	Recruiting
ALL	CD19	Lentivirus	scFv-(4-1BB-CD3 $\zeta$ )	University of Pennsylvania	NCT02030847	Recruiting
ALL	CD19	Lentivirus	scFv-(CD3 $\zeta$ )	Fred Hutchinson Cancer Research Center	NCT01475058	Completed
ALL	CD19	Lentivirus	scFv-(CD3 $\zeta$ )	Fred Hutchinson Cancer Research Center	NCT01865617	Recruiting
ALL	CD19	Lentivirus	scFv-(CD28-CD3 $\zeta$ )	City of Hope Medical Center	NCT02146924	Recruiting
AML	CD123	Lentivirus	scFv-(CD28-CD3 $\zeta$ )	City of Hope Medical Center	NCT02159495	Recruiting
ALL	CD19	Retrovirus	scFv-(CD28-CD3 $\zeta$ & CD28-CD137-CD3 $\zeta$ )	Baylor College of Medicine	NCT01853631	Recruiting
ALL	CD19	Retrovirus	scFv-(CD28-CD3 $\zeta$ )	Memorial Sloan Kettering Cancer Center	NCT01840566	Recruiting
ALL	CD19	Retrovirus/ Lentivirus	scFv-(CD28-CD3 $\zeta$ & 4-1BB-CD3 $\zeta$ )	Memorial Sloan Kettering Cancer Center	NCT00466531	Recruiting
ALL	CD19	Retrovirus	scFv-(CD3 $\zeta$ )	University College London	NCT01195480	Recruiting
ALL	CD19	Retrovirus	scFv-(CD28-CD3 $\zeta$ )	Memorial Sloan Kettering Cancer Center	NCT01860937	Recruiting
ALL	CD19	Retrovirus	scFv-(CD28-CD3 $\zeta$ -4-1BB)	Uppsala University	NCT02132624	Recruiting
ALL	CD19	Lentivirus	scFv-(4-1BB-CD3 $\zeta$ )	Seattle Children's Hospital	NCT02028455	Recruiting
ALL	CD19	Retrovirus	scFv-(CD137-CD3 $\zeta$ )	Chinese PLA General Hospital	NCT01864889	Recruiting
ALL	CD19	Transposon	scFv-(CD3 $\zeta$ )	MD Anderson Cancer Center	NCT01497184	Active, not recruiting
ALL	CD19	Retrovirus	scFv-(CD3 $\zeta$ )	National Cancer Institute	NCT01593696	Recruiting
AML	Lewis-Y	Retrovirus	scFv-(CD28-CD3 $\zeta$ )	Peter MacCallum Cancer Center	NCT01716364	Unknown
ALL	CD19	Retrovirus	scFv-(CD28-CD3 $\zeta$ )	National Cancer Institute	NCT00924326	Recruiting
ALL	CD19	Retrovirus	scFv-(CD28-CD3 $\zeta$ )	Memorial Sloan Kettering Cancer Center	NCT01044069	Recruiting
ALL	CD19	Retrovirus	scFv-(CD3 $\zeta$ )	Memorial Sloan Kettering Cancer Center	NCT01430390	Recruiting
AML	CD33	Retrovirus	scFv-(CD137-CD3 $\zeta$ )	Chinese PLA General Hospital	NCT01864902	Recruiting
ALL	CD19	Lentivirus	scFv-(CD28-CD3 $\zeta$ )	Seattle Children's Hospital	NCT01683279	Recruiting
ALL	CD19	Lentivirus	scFv-(4-1BB-CD3 $\zeta$ )	University of Pennsylvania	NCT01747486	Recruiting
ALL	CD19	Retrovirus	scFv-(CD28-CD3 $\zeta$ )	Baylor College of Medicine	NCT00586391	Active, not recruiting
ALL	CD19	Lentivirus	scFv-(4-1BB-CD3 $\zeta$ )	University of Pennsylvania	NCT01626495	Recruiting
ALL	CD19	Lentivirus	scFv-(4-1BB-CD3 $\zeta$ )	University of Pennsylvania	NCT01029366	Completed
ALL	CD19	Retrovirus	scFv-(CD3 $\zeta$ )	Baylor College of Medicine	NCT00840853	Recruiting
ALL	CD19	Transposon	scFv-(CD3 $\zeta$ )	MD Anderson Cancer Center	NCT01362452	Active, not recruiting

ducted preclinical trials using autologous T cells transduced with CD123-targeted CAR (CD123-CAR-T) in infused AML xenograft model. Their data revealed a CD123-CAR-T-mediated potent activity against CD123<sup>+</sup> cell lines as well as primary AML samples. Additionally, CD123-CAR-T cells exhibited anti-leukemic activity *in vivo* in a xenogeneic AML model. Gill and colleagues suggested that CAR-T123-based myeloablation might be used as a novel conditioning regimen for hematopoietic cell transplantations (Gill et al., 2014).

## DISCUSSION

A series of clinical trials point to key factors, including CAR-T cell design, different gene transfer approaches, and tumor burden, that impact efficacy of the CAR-T cell treatment of hematologic malignancies. Preclinical studies and clinical trials have demonstrated the advantage of CAR-T cell *in vivo* antitumor activity, enhanced by second- and third-generation CAR containing T cell co-stimulatory signaling domains, when compared with first-generation CAR

(Brentjens et al., 2007; Teng et al., 2004). For example, 2014 work of Maude and colleagues (Maude et al., 2014) on second-generation CAR-T cells and ALL treatment, demonstrated superiority of this approach compared with earlier first-generation CAR-T cell experiments. Furthermore, most investigators have been attempting different T cell gene transfer techniques influencing the efficacy of CAR-T cell applications. Retroviruses have been frequently used in ALL research because they efficiently and permanently transduce T cells, and preliminary studies verified safety of this approach (Scholler et al., 2012). Lentiviral vectors also efficiently transduce T cells but are expensive to manufacture, yet potentially safer than retroviruses (Biffi et al., 2011). Investigators have seldom resorted to electroporation or transposon gene transfer technologies (Jin et al., 2011; Till et al., 2008). Alas, no published clinical studies directly compare the different gene transfer approaches, leaving open the question of whether clinically meaningful differences exist between these gene transfer vectors. Tumor burden was demonstrated to be yet another important factor in CAR-T efficacy in a 2011 study by Memorial Sloan-Kettering Cancer Center, where an inverse correlation was noted between tumor burden and the detectable infused modified T cells over time (Brentjens et al., 2011). Most trials focused on the protocols of allo-HSCT in combination with CAR-T cells, superior to CAR-T cell approach alone, until 2014, when Maus and colleagues hypothesized that CAR-T cells may replace allogeneic transplantation as a definitive therapy. Maude has published results of a clinical trial that verified this hypothesis (Maude et al., 2014): the event-free survival and OS of ALL patients who had previously undergone stem-cell transplantation were not different after CAR-T cell treatment than in ALL patients who have not received stem-cell transplantation. Integration of CAR-T with allo-HSCT in future treatment protocols is worth considering, and CAR-T and allo-HSCT timings are expected to be verified by future large sample randomized trials. CRS is also noteworthy. Investigators have shown that CRS was often accompanied by macrophage activation syndrome (MAS), which may be driven in part by high IL-6 levels (Grupp et al., 2013). In the above-mentioned clinical trials, most patients had some degree of CRS. It is not yet clear whether the severity of CRS or MAS is related to antitumor efficacy. In conclusion, CAR-T cell approach has been, undoubtedly, a breakthrough in leukemia treatment. Numerous international trials are currently under way to optimize, such as CAR design, integration protocol optimization, etc.

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